

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING
NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):



1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
7. Other: _____

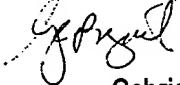
Applicant Must Provide:

- An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g).

For questions regarding compliance to these requirements, please contact:

- For Rules Interpretation, call (703) 308-4216 or (703) 308-2923
- For CRF Submission Help, call (703) 308-4212
- For PatentIn software Program Support:
 - HELP DESK: (703) 739-8559, ext 508, M-F, 8 AM to 5 PM EST except holidays
 - Email: PATIN21HELP@uspto.gov
 - To purchase PatentIn software: (703) 306-2600

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE


 Gabriele E. Bugaisky
 AU 1653
 3/15/01 (703)308-4201

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION, US/09/493,795DATE: 02/17/2000
TIME: 12:51:55

Input Set: I493795.RAW



This Raw Listing contains the General Information
Section and up to first 5 pages.

PP523
Does Not Comply
Corrected Diskette Needed

110> APPLICANT: Watkins, Maren
Olivera, Baldomero M.
Hillyard, David R.
McIntosh, J. Michael
Jones, Robert M.
<120> TITLE OF INVENTION: Alpha-Conotoxin Peptides
<130> FILE REFERENCE: Alphas 2
<140> CURRENT APPLICATION NUMBER: US/09/493,795
<141> CURRENT FILING DATE: 2000-01-28
<150> EARLIER APPLICATION NUMBER: US 60/118,381
<151> EARLIER FILING DATE: 1999-01-29
<160> NUMBER OF SEQ ID NOS: 404
<170> SOFTWARE: PatentIn Ver. 2.0
<210> SEQ ID NO 1
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence:Alpha-Conotoxin
Peptide Generic Formula I
<220> FEATURE:
<221> NAME/KEY: PEPTIDE
<222> LOCATION: (1)...(3)
<223> OTHER INFORMATION: Xaa at residue 1 is des-Xaa, Ile, Leu or Val; Xaa
at residue 2 is des-Xaa, Ala or Gly; Xaa at
residue 3 is des-Xaa, Gly, Trp (D or L), neo-Trp,
halo-Trp or any unnatural aromatic amino acid.
<220> FEATURE:
<221> NAME/KEY: PEPTIDE
<222> LOCATION: (4)...(5)
<223> OTHER INFORMATION: Xaa at residue 4 is des-Xaa, Gly, Trp (D or L),
neo-Trp, halo-Trp or any unnatural aromatic amino
acid; Xaa at residue 5 is Glu, gamma-carboxy-Glu
(Gla), Asp, Ala, Thr, Ser, Gly, Ile, Tyr, nor-Tyr,
<220> FEATURE:
<221> NAME/KEY: PEPTIDE
<222> LOCATION: (5)...(8)
<223> OTHER INFORMATION: mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy
containing amino acid; Xaa at residue 8 is Ser,
Thr, Arg, ornithine, homoarginine, Lys,
N-methyl-Lys,
<220> FEATURE:
<221> NAME/KEY: PEPTIDE

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RAW SEQUENCE LISTING
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45 <222> LOCATION: (8)..(9)
46 <223> OTHER INFORMATION: N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any
47 unnatural basic amino acid; Xaa at residue 9 is
48 Asp, Glu, Gla, Arg, ornithine, homoarginine, Lys,
49 N-methyl-Lys, N,N-dimethyl-Lys,
50 N,N,N-trimethyl-Lys or
51 <220> FEATURE:
52 <221> NAME/KEY: PEPTIDE
53 <222> LOCATION: (9)..(11)
54 <223> OTHER INFORMATION: any unnatural basic amino acid; Xaa at residue 10
55 is Ser, Thr, Asn, Ala, Gly, His, halo-His, Pro or
56 hydroxy-Pro; Xaa at residue 11 is Thr, Ser, Ala,
57 Asp, Asn, Pro, hydroxy-Pro,
58 <220> FEATURE:
59 <221> NAME/KEY: PEPTIDE
60 <222> LOCATION: (11)..(13)
61 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,
62 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any
63 unnatural basic amino acid; Xaa at residue 13 is
64 Gly, Ser, Thr, Ala, Asn,
65 <220> FEATURE:
66 <221> NAME/KEY: PEPTIDE
67 <222> LOCATION: (13)..(14)
68 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,
69 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any
70 unnatural basic amino acid; Xaa at residue 14 is
71 Gln, Leu, His, halo-His, Trp (D or L), halo-Trp,
72 neo-Trp,
73 <220> FEATURE:
74 <221> NAME/KEY: PEPTIDE
75 <222> LOCATION: (14)
76 <223> OTHER INFORMATION: Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
77 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr, Arg,
78 ornithine, homoarginine, Lys, N-methyl-Lys,
79 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any
80 unnatural basic amino
81 <220> FEATURE:
82 <221> NAME/KEY: PEPTIDE
83 <222> LOCATION: (14)..(15)
84 <223> OTHER INFORMATION: acid or any unnatural aromatic amino acid; Xaa at
85 residue 15 is Asn, His, halo-His, Ile, Leu, Val,
86 Gln, Arg, ornithine, homoarginine, Lys,
87 N-methyl-Lys, N,N-dimethyl-Lys,
88 N,N,N-trimethyl-Lys or any
89 <220> FEATURE:
90 <221> NAME/KEY: PEPTIDE
91 <222> LOCATION: (15)..(16)
92 <223> OTHER INFORMATION: unnatural basic amino acid; Xaa at residue 16 is
93 des-Xaa, Val, Ile, Leu, Arg, ornithine,
94 homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,

*Cys is at residue 16 - do you mean
17
Xaa
is at
location*

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RAW SEQUENCE LISTING
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95 N,N,N-trimethyl-Lys or any unnatural basic amino
96 acid.

97 <400> SEQUENCE: 1 Xaa Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Cys 15
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W--> 99 Xaa
100 100
101 <210> SEQ ID NO 2
102 <211> LENGTH: 21
103 <212> TYPE: PRT
104 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: Description of Artificial Sequence:Alpha-Conotoxin
107 Peptide Generic Formula II.
108 <220> FEATURE:
109 <221> NAME/KEY: PEPTIDE
110 <222> LOCATION: (1)..(3)
111 <223> OTHER INFORMATION: Xaa at residue 1 is des-Xaa, Asp, Glu or
112 gamma-carboxy-Glu (Gla); Xaa at residue 2 is
113 des-Xaa, Gln, Ala, Asp, Glu, Gla; Xaa at residue 3
114 is des-Xaa, Gly, Ala, Asp, Glu, Gla, Pro or
115 hydroxy-Pro.
116 <220> FEATURE:
117 <221> NAME/KEY: PEPTIDE
118 <222> LOCATION: (4)..(7)
119 <223> OTHER INFORMATION: Xaa at residue 4 is des-Xaa4, Gly, Glu, Gla, Gln,
120 Asp, Asn, Pro or hydroxy-Pro; Xaa at residue 7 is
121 Ser, Thr, Gly, Glu, Gla, Asn, Trp (D or L),
122 neo-Trp, halo-Trp, Arg, ornithine, homoarginine,
123 <220> FEATURE:
124 <221> NAME/KEY: PEPTIDE
125 <222> LOCATION: (7)
126 <223> OTHER INFORMATION: Lys, N-methyl-Lys, N,N-dimethyl-Lys,
127 N,N,N-trimethyl-Lys, any unnatural basic amino
128 acid, Tyr, nor-Tyr, mono-halo-Tyr, di-halo-Tyr,
129 O-sulpho-Tyr, O-phospho-Tyr, nitro-Tyr or any
130 unnatural hydroxy
131 <220> FEATURE:
132 <221> NAME/KEY: PEPTIDE
133 <222> LOCATION: (7)..(8)
134 <223> OTHER INFORMATION: containing amino acid; Xaa at residue 8 is Asp,
135 Asn, His, halo-His, Thr, Ser, Tyr, nor-Tyr,
136 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
137 O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy
138 <220> FEATURE:
139 <221> NAME/KEY: PEPTIDE
140 <222> LOCATION: (8)..(10)
141 <223> OTHER INFORMATION: containing amino acid; Xaa at residue 9 is Pro or
142 hydroxy-Pro; Xaa at residue 10 is Ala, Ser, Thr,
143 Asp, Val, Ile, Pro, hydroxy-Pro, Tyr, nor-Tyr,
144 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/493,795DATE: 02/17/2000
TIME: 12:51:55

Input Set: I493795.RAW

145 <220> FEATURE:
146 <221> NAME/KEY: PEPTIDE
147 <222> LOCATION: (10)..(12)
148 <223> OTHER INFORMATION: O-phospho-Tyr, nitro-Tyr or any unnatural hydroxy
149 containing amino acid; Xaa at residue 12 is Gly,
150 Ile, Leu, Val, Ala, Thr, Ser, Pro, hydroxy-Pro,
151 Phe, Trp (D or L), neo-Trp, halo-Trp, Arg,
152 ornithine,
153 <220> FEATURE:
154 <221> NAME/KEY: PEPTIDE
155 <222> LOCATION: (12)..(13)
156 <223> OTHER INFORMATION: homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
157 N,N,N-trimethyl-Lys, any unnatural basic amino
158 acid or any unnatural aromatic amino acid; Xaa at
159 residue 13 is Ala, Asn, Phe, Pro, hydroxy-Pro,
160 <220> FEATURE:
161 <221> NAME/KEY: PEPTIDE
162 <222> LOCATION: (13)
163 <223> OTHER INFORMATION: Glu, Gla, Gln, His, halo-His, Val, Ser, Thr, Arg,
164 ornithine, homoarginine, Lys, N-methyl-Lys,
165 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys or any
166 unnatural basic amino acid.
167 <220> FEATURE:
168 <221> NAME/KEY: PEPTIDE
169 <222> LOCATION: (14)
170 <223> OTHER INFORMATION: Xaa at residue 14 is Thr, Ser, His, halo-His, Leu,
171 Ile, Val, Asn, Met, Pro, hydroxy-Pro, Arg,
172 ornithine, homoarginine, Lys, N-methyl-Lys,
173 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any
174 unnatural basic
175 <220> FEATURE:
176 <221> NAME/KEY: PEPTIDE
177 <222> LOCATION: (14)..(15)
178 <223> OTHER INFORMATION: amino acid, Tyr, nor-Tyr, mono-halo-Tyr,
179 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,
180 nitro-Tyr or any unnatural hydroxy containing
181 amino acid; Xaa at residue 15 is Asn, Pro,
182 hydroxy-Pro, Gln, Ser, Thr,
183 <220> FEATURE:
184 <221> NAME/KEY: PEPTIDE
185 <222> LOCATION: (15)
186 <223> OTHER INFORMATION: Arg, ornithine, homoarginine, Lys, N-methyl-Lys,
187 N,N-dimethyl-Lys N,N,N-trimethyl-Lys, any
188 unnatural basic amino acid, Tyr, nor-Tyr,
189 mono-halo-Tyr, di-halo-Tyr, O-sulpho-Tyr,
190 O-phospho-Tyr, nitro-Tyr
191 <220> FEATURE:
192 <221> NAME/KEY: PEPTIDE
193 <222> LOCATION: (15)..(16)
194 <223> OTHER INFORMATION: or any unnatural hydroxy containing amino acid;

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RAW SEQUENCE LISTING
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195 Xaa at residue 16 is des-Xaa, Gly, Thr, Ser, Pro,
 196 hydroxy-Pro, Tyr, nor-Tyr, mono-halo-Tyr,
 197 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,
 198 nitro-Tyr or any
 199 <220> FEATURE:
 200 <221> NAME/KEY: PEPTIDE
 201 <222> LOCATION: (16)..(17)
 202 <223> OTHER INFORMATION: unnatural hydroxy containing amino acid; Xaa at
 203 residue 17 is des-Xaa14, Ile, Val, Asp, Leu, Phe,
 204 Arg, ornithine, homoarginine, Lys, N-methyl-Lys,
 205 N,N-dimethyl-Lys, N,N,N-trimethyl-Lys, any
 206 unnatural
 207 <220> FEATURE:
 208 <221> NAME/KEY: PEPTIDE
 209 <222> LOCATION: (17)..(19)
 210 <223> OTHER INFORMATION: basic amino acid, Tyr, nor-Tyr, mono-halo-Tyr,
 211 di-halo-Tyr, O-sulpho-Tyr, O-phospho-Tyr,
 212 nitro-Tyr or any unnatural hydroxy containing
 213 amino acid; Xaa at residue 19 is des-Xaa, Gly,
 214 Ala, Met, Ser,
 215 <220> FEATURE:
 216 <221> NAME/KEY: PEPTIDE
 217 <222> LOCATION: (19)
 218 <223> OTHER INFORMATION: Thr, Trp (D or L), neo-Trp, halo-Trp, any
 219 unnatural aromatic amino acid, Arg, ornithine,
 220 homoarginine, Lys, N-methyl-Lys, N,N-dimethyl-Lys,
 221 N,N,N-trimethyl-Lys or any unnatural basic amino
 222 acid.
 223 <220> FEATURE:
 224 <221> NAME/KEY: PEPTIDE
 225 <222> LOCATION: (20)
 226 <223> OTHER INFORMATION: Xaa at residue 20 is des-Xaa, Trp (D or L),
 227 neo-Trp, halo-Trp, any unnatural aromatic amino
 228 acid, Arg, ornithine, homoarginine, Lys,
 229 N-methyl-Lys, N,N-dimethyl-Lys,
 230 N,N,N-trimethyl-Lys or any
 231 <220> FEATURE:
 232 <221> NAME/KEY: PEPTIDE
 233 <222> LOCATION: (20)..(21)
 234 <223> OTHER INFORMATION: unnatural basic amino acid; Xaa at residue 21 is
 235 des-Xaa, Arg, ornithine, homoarginine, Lys,
 236 N-methyl-Lys, N,N-dimethyl-Lys,
 237 N,N,N-trimethyl-Lys or any unnatural basic amino
 238 acid.
 239 <400> SEQUENCE: 2
 240 Xaa Xaa Xaa Xaa Cys Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa
 241 1 5 10 15
 242 Xaa Cys Xaa Xaa Xaa
 243 20

Please Note: <210> SEQ ID NO 3

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
 Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
 <223> fields of each sequence which presents at least one n or Xaa.

PJI

Input Set: I493795.RAW

Line	?	Error/Warning	Original Text
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242	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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642	W	"N" or "Xaa" used: Feature required	Ala Xaa Gly Cys Cys Asn Xaa Ala Cys V
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693	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Gly Asn Xaa Asp Cys Thr Ser H
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890	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ser Gly A
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903	W	"N" or "Xaa" used: Feature required	Xaa Cys Cys Ser His Xaa Ala Cys Ser Gly A
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1037	W	"N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn S

Input Set: I493795.RAW

Line ? Error/Warning

Original Text

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 1076 W "N" or "Xaa" used: Feature required
 1088 W "N" or "Xaa" used: Feature required
 1108 W "N" or "Xaa" used: Feature required
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 1129 W "N" or "Xaa" used: Feature required
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 2895 W "N" or "Xaa" used: Feature required
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 2948 W "N" or "Xaa" used: Feature required
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 2987 W "N" or "Xaa" used: Feature required
 2999 W "N" or "Xaa" used: Feature required
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 3032 W "N" or "Xaa" used: Feature required
 3046 W "N" or "Xaa" used: Feature required
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 Xaa Val Cys Cys Ser Asp Xaa Arg Cys Asn V
 Gly Cys Cys Ser Arg Xaa Xaa Cys Ile Ala A
 Xaa Gln Cys Cys Ser His Leu Ala Cys Asn V
 Gly Cys Cys Ser Xaa Phe Asp Cys Arg Met M
 Gly Xaa Arg
 Gly Gly Cys Cys Ser Phe Ala Ala Cys Arg X
 Gly Gly Cys Cys Phe His Xaa Val Cys Xaa I
 Ser Ala Thr Cys Cys Asn Xaa Xaa Cys X
 Ala Cys Cys Ala Xaa Xaa Xaa Cys Phe Xaa A
 Asn Ala Xaa Cys Cys Xaa Xaa Xaa Cys X
 Gly Cys Cys Xaa Asn Xaa Val Cys Xaa Xaa X
 Xaa Xaa Xaa Gly Cys Cys Arg His Xaa Ala C
 Cys Cys Ala Asp Xaa Asp Cys Arg Phe Arg X
 Gly Cys Cys Xaa Asn Xaa Ser Cys Xaa Xaa X
 Xaa
 Cys Cys Ser Asn Xaa Thr Cys Xaa Xaa Thr X
 Cys Cys Ala Asn Xaa Ile Cys Xaa Asn Thr X
 Cys Cys Asn Asn Xaa Thr Cys Xaa Xaa Thr X
 Cys Cys Ser Asn Xaa Val Cys Xaa Xaa Thr X
 Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile A
 Gly Cys Cys Ser His Xaa Val Cys Ser Ala M
 Gly Cys Cys Xaa Asn Xaa Xaa Cys Gly Ala S
 Gly Cys Cys Ser Xaa Xaa Xaa Cys Phe Ala T
 Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile A
 Gly Gly Cys Cys Ser Xaa Xaa Xaa Cys Ile A
 Asp Cys Cys Ser Asn Xaa Xaa Cys Ser Gln A
 Asp Cys Cys Ser Asn Xaa Xaa Cys Ala His A
 Xaa Cys Cys Thr Asn Xaa Val Cys His Ala X
 Gly Cys Cys Ser Asn Xaa Val Cys His Leu X
 Xaa Cys Cys Thr Asn Xaa Val Cys His Val X
 Xaa Xaa Cys Cys Ser Xaa Xaa Ala Cys Asn L
 Xaa Xaa Cys Cys Ser Asp Xaa Arg Cys Asn S
 Leu Asn Cys Cys Met Ile Xaa Xaa Cys Xaa X
 Cys Ser Xaa Val Arg
 Ala Phe Gly Cys Cys Asp Leu Ile Xaa Cys L
 Arg Cys Asn Xaa Val His
 Leu Gly Cys Cys Asn Val Thr Xaa Cys Xaa X
 Cys Asn Xaa Val Arg
 Asp Xaa Cys Cys Ser Asn Xaa Ala Cys Arg V
 Leu Asn Cys Cys Ser Ile Xaa Gly Cys Xaa A
 Cys Ser Xaa Val Arg
 Gly Gly Cys Cys Ser His Xaa Val Cys Xaa P
 Gly Gly Cys Cys Ser His Xaa Val Cys Asn L
 Gly Cys Cys Ser His Xaa Xaa Cys Xaa Ala A
 Gly Gly Cys Cys Ser His Xaa Ala Cys Ser V
 Gly Gly Cys Cys Ser Xaa Xaa Ala Cys Ser V
 Val Ser Cys Cys Val Val Arg Xaa Cys Xaa I
 Cys Leu Xaa Ala Asp Xaa Arg Thr Leu
 Xaa Asn Cys Cys Ser Ile Xaa Gly Cys Xaa X

Input Set: I493795.RAW

Line	?	Error/Warning	Original Text
3376	W	"N" or "Xaa" used: Feature required	Cys Ser Xaa Val Arg
3388	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys His Leu X
3402	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Ile Cys Xaa Phe A
3422	W	"N" or "Xaa" used: Feature required	Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa X
3434	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly A
3445	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ala Val Xaa Ser Cys Arg Leu A
3457	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Asn Val A
3470	W	"N" or "Xaa" used: Feature required	Thr Xaa Xaa Xaa Cys Xaa Asn Xaa Xaa C
3485	W	"N" or "Xaa" used: Feature required	Asp Ala Cys Cys Ser Asp Xaa Arg Cys Ser G
3498	W	"N" or "Xaa" used: Feature required	Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser V
3510	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly S
3522	W	"N" or "Xaa" used: Feature required	Xaa Asp Cys Cys Ser Asp Xaa Arg Cys Ser V
3534	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Ala Gly A
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3561	W	"N" or "Xaa" used: Feature required	Thr Asp Ser Xaa Xaa Cys Cys Leu Asp Ser A
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3608	W	"N" or "Xaa" used: Feature required	Gly Xaa Cys Cys Ile Asn Asp Ala Cys Arg S
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3679	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Val Cys Xaa Gln A
3681	W	"N" or "Xaa" used: Feature required	Arg Xaa Ser
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3725	W	"N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Xaa Xaa Asp Cys Gly A
3745	W	"N" or "Xaa" used: Feature required	Xaa Cys Cys Ser Gln Xaa Xaa Cys Arg Xaa X
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3771	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asp Xaa Ser Cys Asn Val A
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3797	W	"N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser Asn Xaa Ala Cys Leu V
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3822	W	"N" or "Xaa" used: Feature required	Cys Cys Ser Asp Xaa Ser Cys Xaa Arg Leu H
3835	W	"N" or "Xaa" used: Feature required	Cys Cys Thr Asn Xaa Ala Cys Leu Val Asn A
3847	W	"N" or "Xaa" used: Feature required	Asp Xaa Cys Cys Ser Asp Xaa Arg Cys His G
3859	W	"N" or "Xaa" used: Feature required	Asp Cys Cys Ser His Xaa Leu Cys Arg Leu P
3879	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Xaa Val A
3891	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Ala Cys Asn Val A
3910	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Xaa Val A
3923	W	"N" or "Xaa" used: Feature required	Gly Gly Cys Cys Ser His Xaa Ala Cys Xaa V
3936	W	"N" or "Xaa" used: Feature required	Val Cys Cys Ser Asn Xaa Val Cys His Val A
3949	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser His Xaa Val Cys Asn Leu S
3963	W	"N" or "Xaa" used: Feature required	Xaa Xaa Cys Cys Ser His Xaa Ala Cys Asn V
3975	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Ser Asn Xaa Ala Cys Leu Val A
3987	W	"N" or "Xaa" used: Feature required	Asp Cys Cys Asp Asp Xaa Ala Cys Thr Val A
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4024	W	"N" or "Xaa" used: Feature required	Cys Cys Thr Ile Xaa Ser Cys Xaa Xaa Xaa X
4038	W	"N" or "Xaa" used: Feature required	Gly Cys Cys Gly Asn Xaa Ala Cys Ser Gly S
4357	W	Invalid/Missing Amino Acid Numbering	

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VERIFICATION SUMMARY
PATENT APPLICATION US/09/493,795

DATE: 02/17/2000
TIME: 12:51:55

Input Set: I493795.RAW

Line ? Error/Warning

Original Text

4440 W Invalid/Missing Amino Acid Numbering